V3 loop sequence data of HIV-1 patient isolates (PI)

CTRPNNNTRKSI.HIGPGRAFYATGDIIGDIRQAHC

TODITEDIROAHO
GSTNAS
HNW-T
SEE
EE
EE
-I
-IH-TVTDRS-HT-RKIK
SIQK-R-V.RS-I-ARAATK-Q-
SIQK-R-V.RS-I-RAATK-Q- YR-AKHR-MK-Q-
YR-AKHR-MS-IRAATK-Q-

Amino acid sequences of the NL4-3 V2 loop and NL4-3 V3 loop. The regions which are preferably to be varied are underlined.

NL4-3 V2 I oop

GEIKNCSFN<u>ISTSIRDKVQKEY</u>AFFYKL<u>DIVPIDNTSY</u>RLISCNTSVITQA

NL4-3 V3 I oop

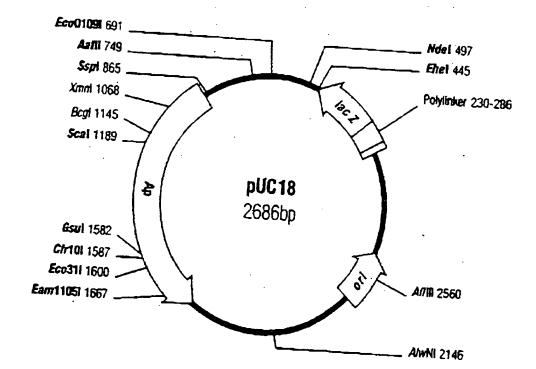
VQL<u>NTS</u>VEI<u>N</u>CTRP<u>NNNTRKSIRIOR</u>GPG<u>RAFVTIGKIGNM</u>RQAHC<u>NIS</u>RAKW<u>NAT</u>LKQ

Variation of the V3 loop. Data from the Los Alamos

ス^ペ・トーにらほれぐされ ٢٠٠٠ م ヹ゚ゝ゚゚゚゚゚゙゙ゔゖ゙ヹヹ゙゙゙゙゙゙゙゙゙゙゙゙゚ 4 5. 0. F. ೦ೄಸ್ಮಪ್ರ≖. ಳ ∝್ಜಸ್.ಬ. ೦, F. ಸ. -gF、シロスペス・ Og Z m よらなよいれっぺいとっ ۵ پاران کا جر <u>゠゚ゝ゚゚゚゚ゖ゚゙゙゙゙゙゙゙゙゙゙゙゙゙゙゚゙゙ゖ゚ヹ゚゚゙゙゙゚ゔ゚゚゚゚゚゚ヸ゚ヸ゚ヹ゚゚゙゙</u>゚ゔ゚゚゚ -%ン'- モリスススペ ·ਸ਼ੵੑੑੑੑੑੑੑਲ਼ੵੑਫ਼ੑਲ਼ੑਫ਼ੑਫ਼ੑਫ਼ੑਫ਼ੑਫ਼੶ਜ਼ਜ਼੶ਲ਼ਜ਼ਜ਼ਫ਼ੑਖ਼ਖ਼ 0 . m x U X 4 F 0 % F. 4. 0. 5. 2. 2. 2. 3. 7. ゖ゚ヺヿ゙゚ゞ゠゙゙゙゙゙゙゙゙゙゙゙ゕ゚ヹ゙゙゙゙゙゙゙゙゙゙゙゙゙ヹヸ ₹_S≻Forkortoriz; ಸ್ಕರ್ಷ ಬ್ರಂ ಸಹವನ ດູ∝.× ຫ.≯. で。>。つ。レ。。。。。ヾヹゝ゚ヸ. Q < E & OF -,7<u>,</u>2,>r.t.o,7,5,7, ಬ್ರ_{ಕ್ಕ}ಸ್ ರ ೧, ಸ್. ಕ. >, ಸ್. x x o v F x m n o -∝್ಲಿಸ⊼⊼ಎಸ್ಎ್ಳೆಗೆ∺ಪ FgX_-_x,>,<,0, z_z≻_x⊢_=ヸ゚ல゚ヸ゚゙゙ゖ゙゚゙゚゚゚ **たったりしゅんけん** ZSOCIETYOR ≈_g ∾, ⊢, >, r₈=_> ≥ ≥ = 1 ₹ ≥ たっぱらずらは ニュン

Fig. 2b

ζ,



Schematic representation of the process for the preparation of the mixture of gp 120-expressing plasmid vectors.

- Preparation of the degenerated DNA fragments for e.g. the V3 loop
 - a.) Synthesis of single-stranded DNA
 - b.) Hybridization of two complementary oligonucleotides.
- 2.) Cloning of the V3 loop DNA fragments into pUC18 delta env
- 3.) Cloning of the env gene into the gp 120 expression vector pBSCenvATG

V3 loop fragments with degenerated sequence e.g.:

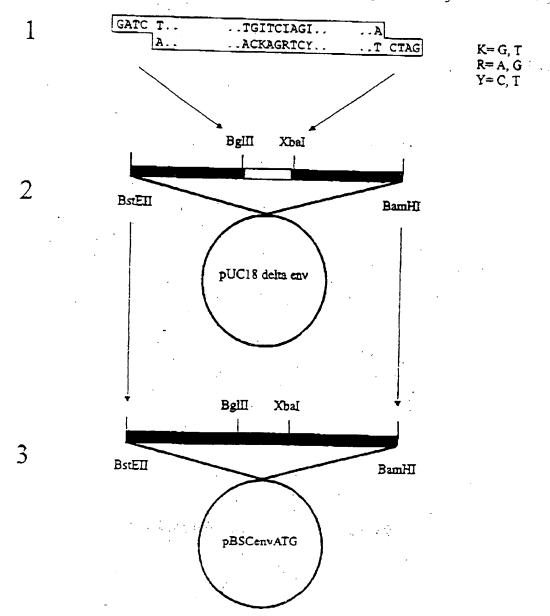


Fig. 5

Heterogeneity of the vaccine using the V3 loop as an example

V3 loop sequence:

CTRPNNNTRKSIHIGPGRAFYTTGDIIGDIRQAHC

GP HAKK R RN ET E

Degeneration at protein level:

 $3 \times 2 \times 2 \times 2 \times 4 \times 6 \times 2 = 1152$ variants

Ser His Tyr Thr Gly Asp Asp
Gly Pro His Ala Lys Lys Glu
Arg Arg Asn
Glu Thr
Glu
Ala

Degenerated DNA sequence of the respective variable amino acid positions:

AGA CCT TAT ACT GGG GAG GAC G T A C G AA ACC G = 2048 variants

AGA CCT TAT ACT GGG GAG GAC
AGT CAT CAT GCT GAG GAC GAG
GGT AGG GCC
AAG
AAC
ACC
ACC

Variability at protein level derived from the degenerated DNA sequence:

Arg Pro Tyr Thr Gly Glu Asp
Ser His His Ala Glu Asp Glu
Gly
Arg Ala
Lys Ala
Lys
Asn
Thr